



# ***Tools for Documenting the Establishment of Microbial Biocontrol Agents***

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**Microbial Biocontrol of Arthropods, Weeds, and Plant  
Pathogens: Risks, Benefits and Challenges**  
Nov. 28-Dec. 1, 2010, National Conservation Training  
Center, Shepherdstown, WV



# Biological Control of Plant Diseases

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## Approaches

- Indigenous organisms: Natural microbially-based defense

### Suppressive Soils

pathogen does not establish or persist

- establishes but causes little or no damage
- establishes, causes disease but then declines in severity
- pathogen may persist in the soil (Baker and Cook, 1974)

- Introduced microbial agents

Many organisms have been studied for biocontrol activity but research has focused on *Pseudomonas*, *Bacillus* & *Trichoderma*.

# Characteristics of Biocontrol Agents

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- *Bacillus*

- ◆ bacterium of choice for commercial BCAs
- ◆ easily mass produced
- ◆ easily formulated; forms a dormant, resistant spore

- *Trichoderma*

- ◆ fungus of choice for commercial BCAs
- ◆ easily mass produced
- ◆ sporulates profusely, easily formulated
- ◆ resistant to natural and synthetic chemicals

- *Pseudomonas*

- ◆ organism of choice for fundamental studies of biocontrol
- ◆ easily mass produced, but harder to formulate; no spores
- ◆ few commercial BCAs

# Tracking Biocontrol Agents



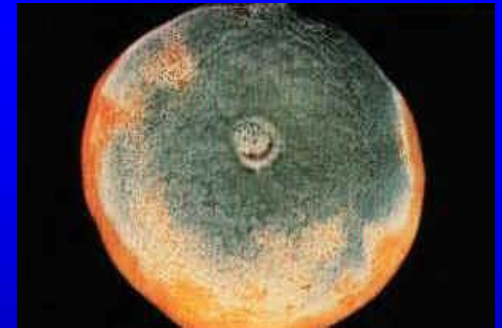
**Foliar diseases**



**Soilborne diseases**



**Postharvest diseases**



**Soil and rhizosphere: most complex environments in which BCAs must establish and function**

**Tracking BCAs: complicated by the microbial milieu in the rhizosphere**



# Characteristics of Root Colonization by Bacterial BCAs

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- BCAs applied to seeds or planting material spread throughout a root system but cells are not uniformly distributed along roots (Bahme & Schroth, 1987; Bull et al., 1991; Loper et al., 1984).
- Densities of BCAs are greatest after planting, decline throughout the growing season or decline & stabilize; populations do not exceed  $5 \times 10^8$  CFU g<sup>-1</sup> root + rhizosphere soil (Bakker et al., 1986; Kluepfel, 1993; Mahaffee et al., 1997; Steddom et al., 2002; Weller, 1983).
- The proportion of the heterotrophic bacterial population comprised by a BCA is greatest soon after planting and decreases throughout the growing season (Halverson et al., 1993; Juhnke et al., 1989; Weller, 1983).

# Tools for Tracking BCAs

## Seeking Sensitivity and Specificity

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- **Direct Microscopy:** (lacks specificity)
  - ◆ **Traditional SEM:** extensive sample preparation
    - Sugar beet seed colonization *B. subtilis* & *P. putida* (Fukui et al., 1994)
    - Distribution of *P. fluorescens* on pea roots (Dandurand et al., 1997)
  - ◆ **Environmental Scanning Electron Microscopy**
    - Water remains liquid and specimens are hydrated; no preparation
- **Molecular Stains:** staining and tagging systems & advanced fluorescence microscopy; detection of single cells in complex environments; greater sensitivity
  - ◆ **General Cell Stains**
    - Epifluorescence microscopy & DNA staining: acridine orange (DeLeo et al., 1997); SYBR Green II (Weinbauer et al., 1998)
    - Fluorescent Brighteners: staining of fungi (Thrane et al., 1999; Eickhorst & Tippkötter, 2008; Harris et al., 2002)

# Tools for Tracking BCAs

## Seeking Sensitivity and Specificity

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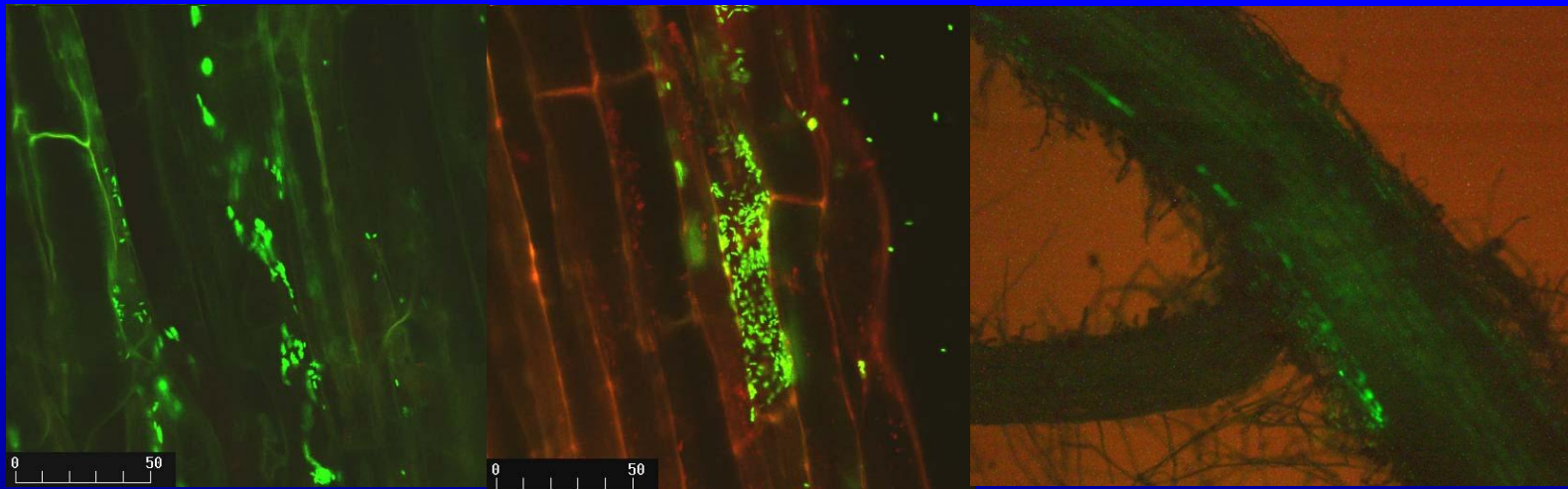
### ◆ Specific Cell Stains

- **Confocal Laser Scanning Microscopy (CLSM) with Strain-Specific Fluorescent Antibody Staining (enhanced sensitivity and specificity)**
  - High resolution 3-D images of structure and composition of microbial communities
  - Early studies: *A. brasilense* and *P. fluorescens* colonizing wheat/ barley roots (Schloter et al., 1993; Hansen et al., 1997; Kirchhof et al., 1997)
- **CLSM with Fluorescence In Situ Hybridization (FISH) (targeting rRNA & mRNA)**
  - Numerous studies of root colonization (Assmus et al., 1995, 1997; Watt et al., 2006; Santaella et al., 2008)
  - FISH showed different colonization patterns for *Salmonella enterica* and *Listeria* spp. on barley roots (Kutter et al., 2005)

# Tools for Tracking BCAs

## Seeking Sensitivity and Specificity

- ◆ **Reporter Strains:** single strains carrying fluorescence marker genes
  - **Non-Specific reporter:** *lux* or *gfp* genes controlled by a constitutive promoter (bioluminescence/ green fluorescent protein)
    - CLSM used to track GFP mutants of *P. chlororaphis* on barley seed (Tombolini et al., 1999); *P. fluorescens* on roots of barley (Normander et al., 1999), tomato (Gotz et al., 2006), avocado (Pliego et al., 2008), olive (Prieto & Mercado-Blanco, 2008), tomato (Chin-A-Woeng et al., 1997; Bloemberg et al., 2000)



CFLM

CFLM

EFM

**Wheat roots colonized by *gfp*-tagged *P. fluorescens* Q8r1-96 (10 days PI)**



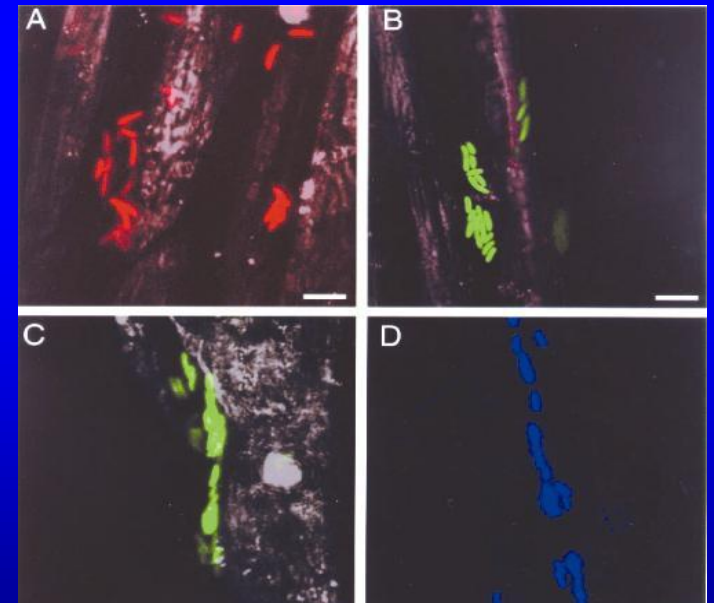
# Tools for Tracking

Simultaneous imaging of *Pseudomonas fluorescens* WCS365 expressing autofluorescent proteins in the rhizosphere.

- GFP - green fluorescent protein (isolated from *Aequorea victoria*)

## Derivatives of GFP

- ECFP - enhanced cyan fluorescent protein; expressing bacteria appear red (A)
- EGFP - enhanced green fluorescent protein; expressing bacteria appear green (B)
- EYFP - enhanced yellow fluorescent protein; expressing bacteria appear green (C)
- RFP - red fluorescent protein; expressing bacteria appear blue (D) (isolated from *Discosoma* spp.)

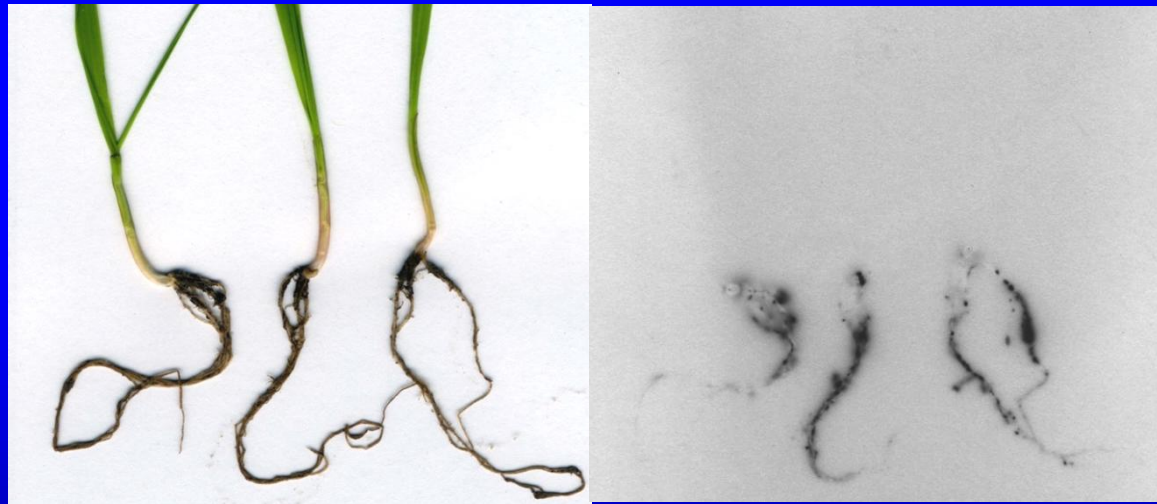


Adapted from Bloembergen et al., 2000

# Tools for Tracking BCAs

## ◆ Reporter Strains:

- **Semi-Specific reporter:** responds to exposure to particular conditions (high temperature; ROS etc.)
- **Specific reporter:** responds to presence (Jager et al., 1999) or absence (Koch et al., 2001) of specific compounds or elements

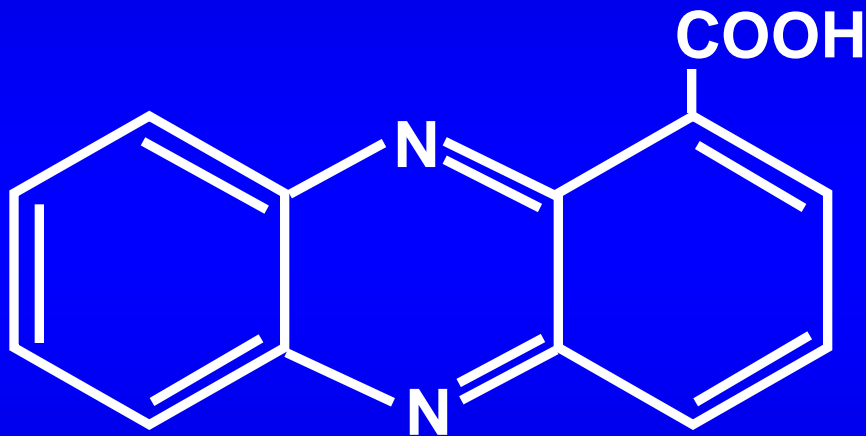


## T3SS genes of Q8r1-96 expressed in the wheat rhizosphere

X-ray film image of roots colonized by Q8r1-96 tagged by a mini-Tn7 transposon with the *rspJ* promoter fused to luciferase genes (Mavrodi et al., in press)

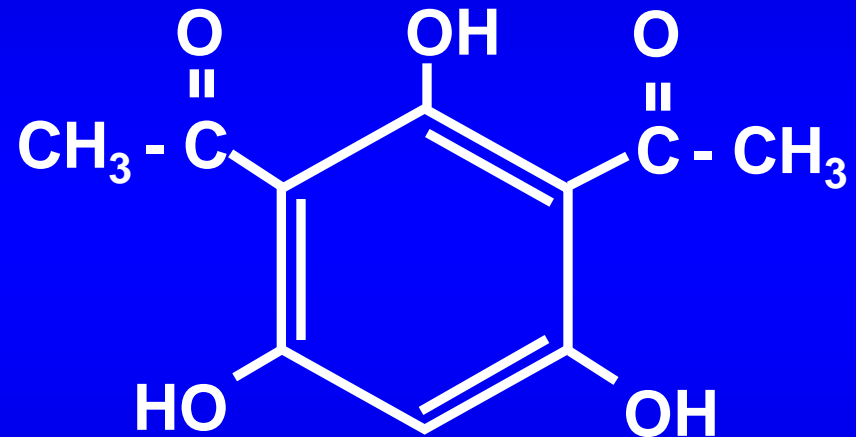
# Well-Studied Groups of BCAs: PCA- & DAPG-Producing *Pseudomonas* spp.

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Phenazine-1-carboxylic acid  
(PCA)

*P. fluorescens* 2-79



2,4-Diacetylphloroglucinol  
(DAPG)

*P. fluorescens* Q8r1-96

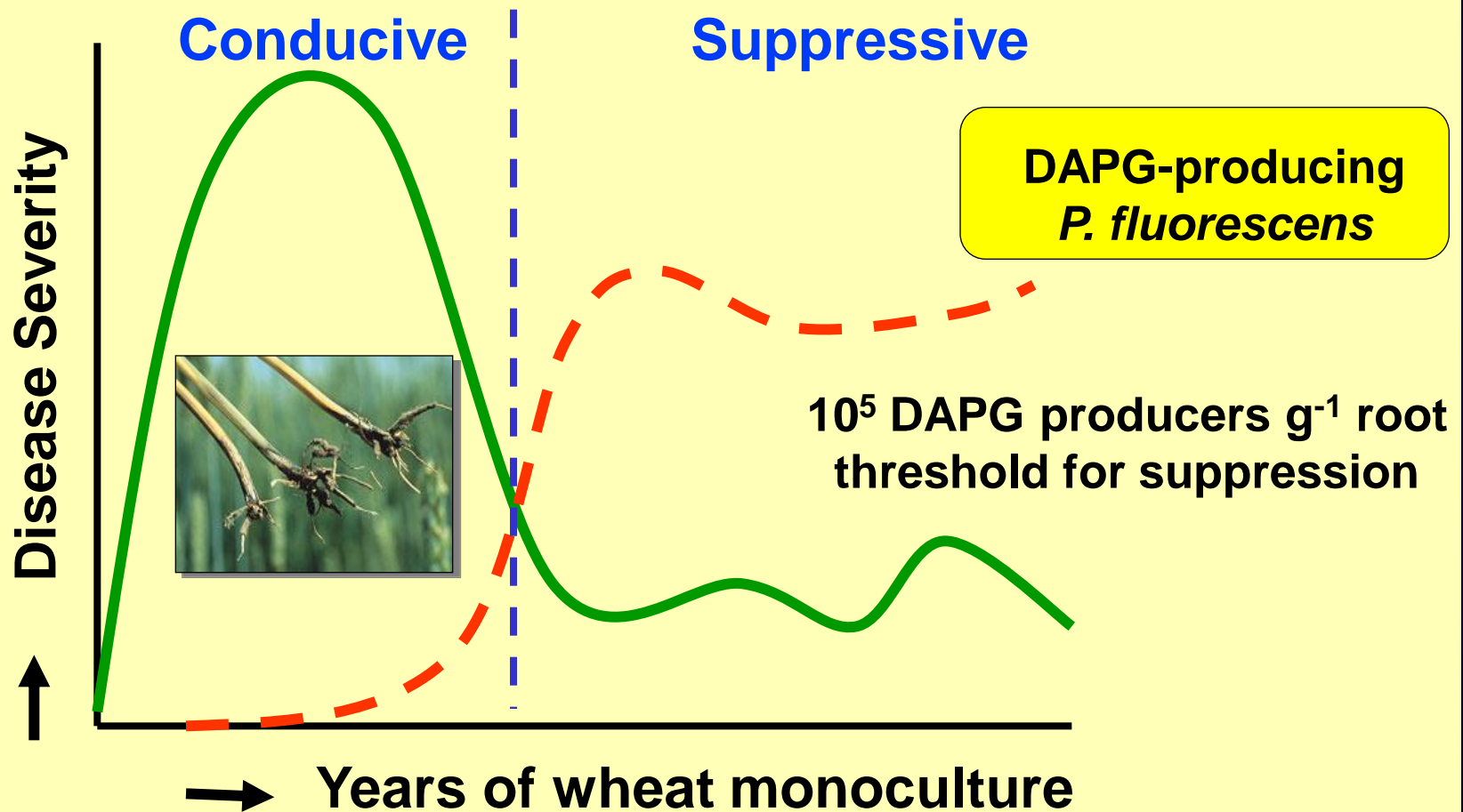
# Take-all

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*Gaeumannomyces  
graminis* var. *tritici*



# Take-All Decline (TAD)





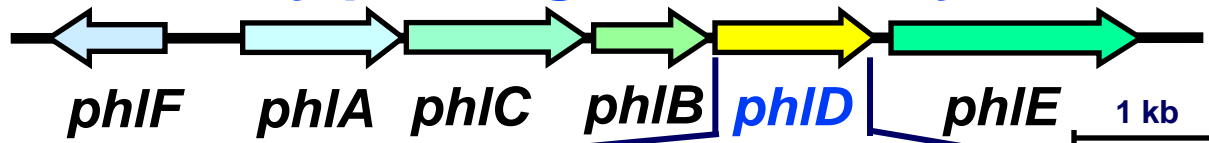
# Detection of DAPG-Producing Fluorescent *Pseudomonas* spp. (*phlD*<sup>+</sup>)

*phlD* encodes for a polyketide synthase; **genetic marker**

*phlD* specific probes and primers used in combination with colony hybridization and PCR to detect DAPG producers

All DAPG producers contain *phlD*

## 2,4-diacetylphloroglucinol biosynthesis genes



**B2BF**

5'-ACCCACCGCAGCATCGTGTATGAGC-3'

**PCR**

629 bp

3'-CCGCCGCTATGGAAGATGAAAAAGTC-5'

**BPR4**

(Bangera & Thomashow. 1999. *J. Bacteriol.*)

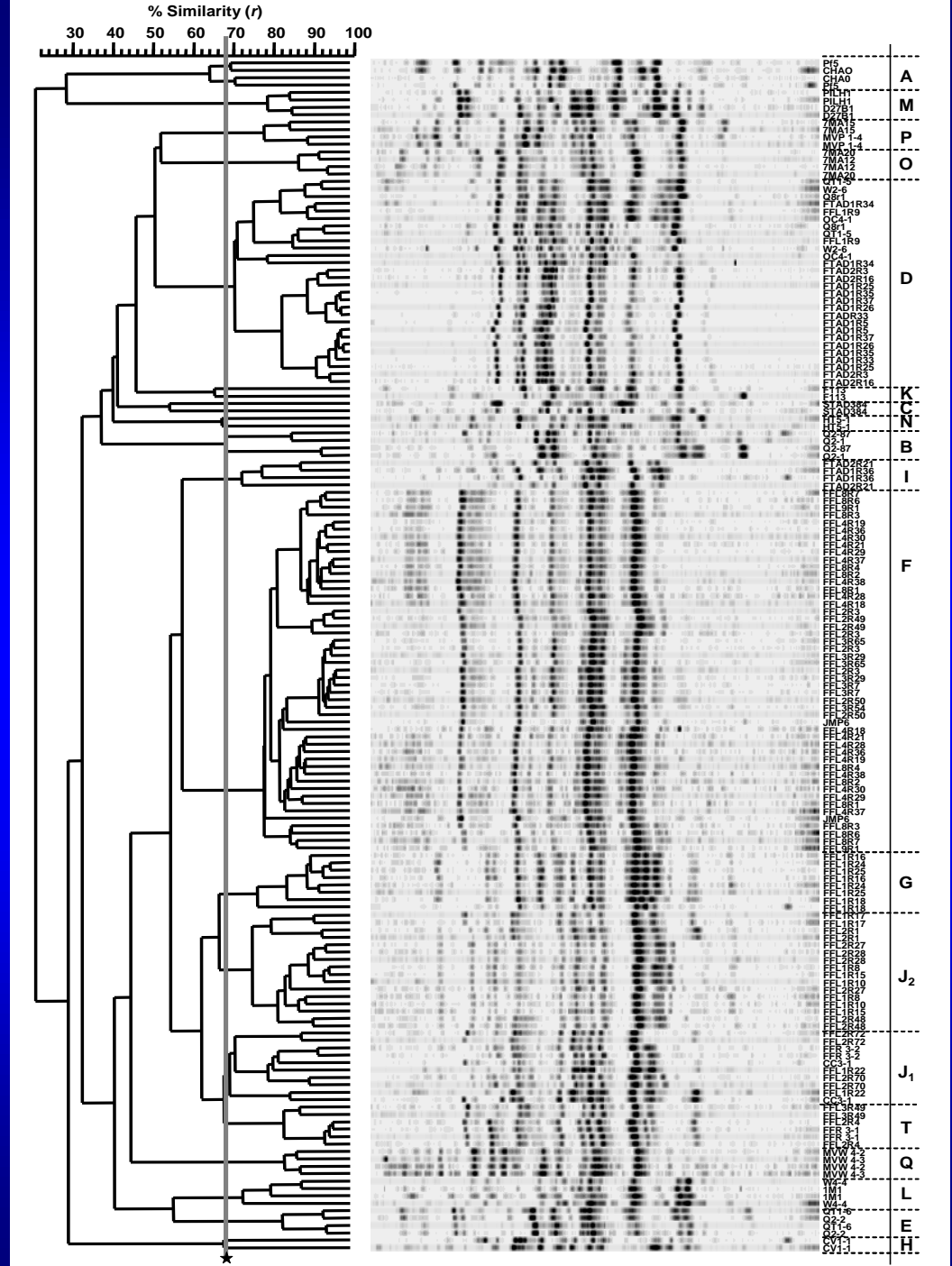
(Raaijmakers et al. 1997. *Appl. Environ. Microbiol.*)

(McSpadden-Gardener et al. 2001. *Phytopathology*)

# Diversity of DAPG Producers

- *phlD*-RFLP
- Rep-PCR (BOXA1R primer)
- 22 Genotypes (A-T, PfY, PfZ)
- D-genotype dominant in Washington TAD soils

Landa et al., 2005  
 Raaijmakers & Weller, 2001  
 Landa et al., 2002  
 McSpadden Gardener et al., 2000



# Tools for Tracking BCAs

## Seeking Sensitivity and Specificity

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- **Culture-Based Methods**

- Selective or Semi-Selective Media**

- Pseudomonas* (Sands & Rovira, 1970; Simon & Ridge, 1974; Grant & Holt, 1977)

- ♦ Antibiotic Resistant Strains**

- Most common approach; rifampicin resistant derivatives (Weller, 1984; Mahaffee et al., 1997)

- ♦ Immunofluorescent Colony Staining (antigenic specificity)**

- Uses strain-specific antibodies; cells remain viable; highly sensitive (Mahaffee et al., 1997; Van der Wolf et al., 1995, 2000)

- ♦ Colony Hybridization (gene-based specificity)**

- Bacterial colonies are replica-plated onto membranes and their DNA is hybridized with a gene-specific probe
    - Identification of DAPG-producing *P. fluorescens* (Raaijmakers et al., 1997)

# Tools for Tracking BCAs

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- Culture-Based Methods:

- ◆ PCR-Based Dilution-Endpoint Method:

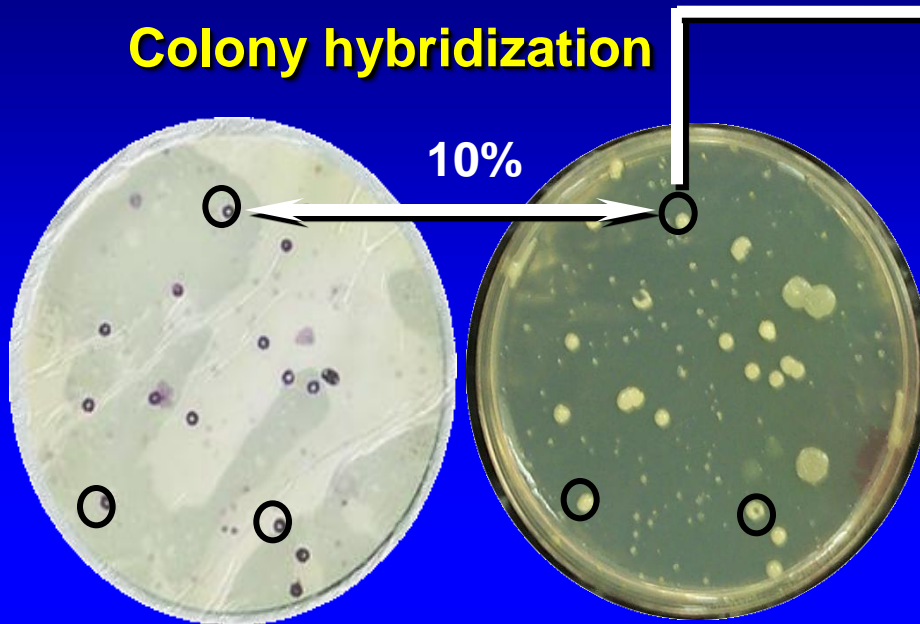
- three fold dilution in microtiter plates with selective media followed by PCR with specific probes (McSpadden Gardener et al., 2001; Landa et al., 2002)

- Culture-Independent Quantitative PCR: (Rezzonico et al., 2003)

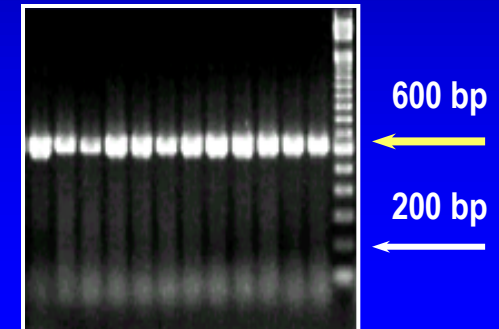
- Mavrodi et al., (2007) used a real-time PCR SYBR green assay to quantify different genotypes of DAPG-producing *P. fluorescens* on the roots of wheat.

# Identification of DAPG-producing *Pseudomonas* spp. from the rhizosphere

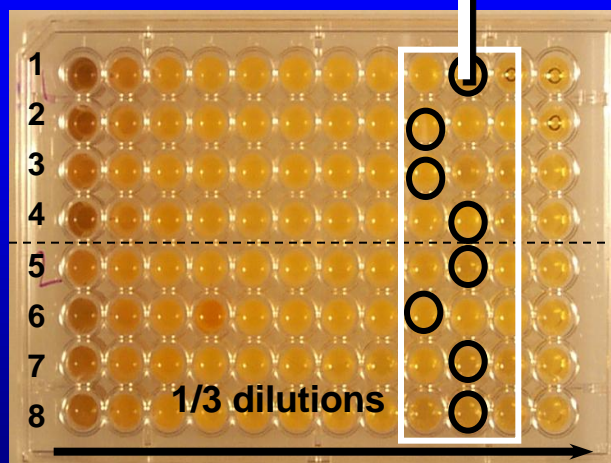
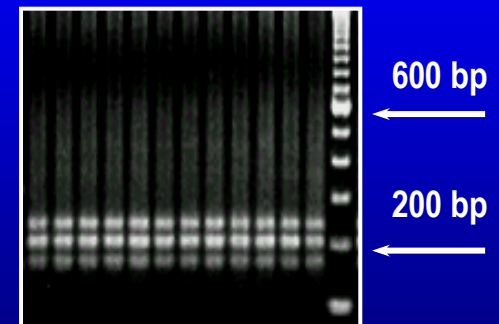
## Colony hybridization



## PCR (B2BF/BPR4)



## RFLP (*Hae* III)



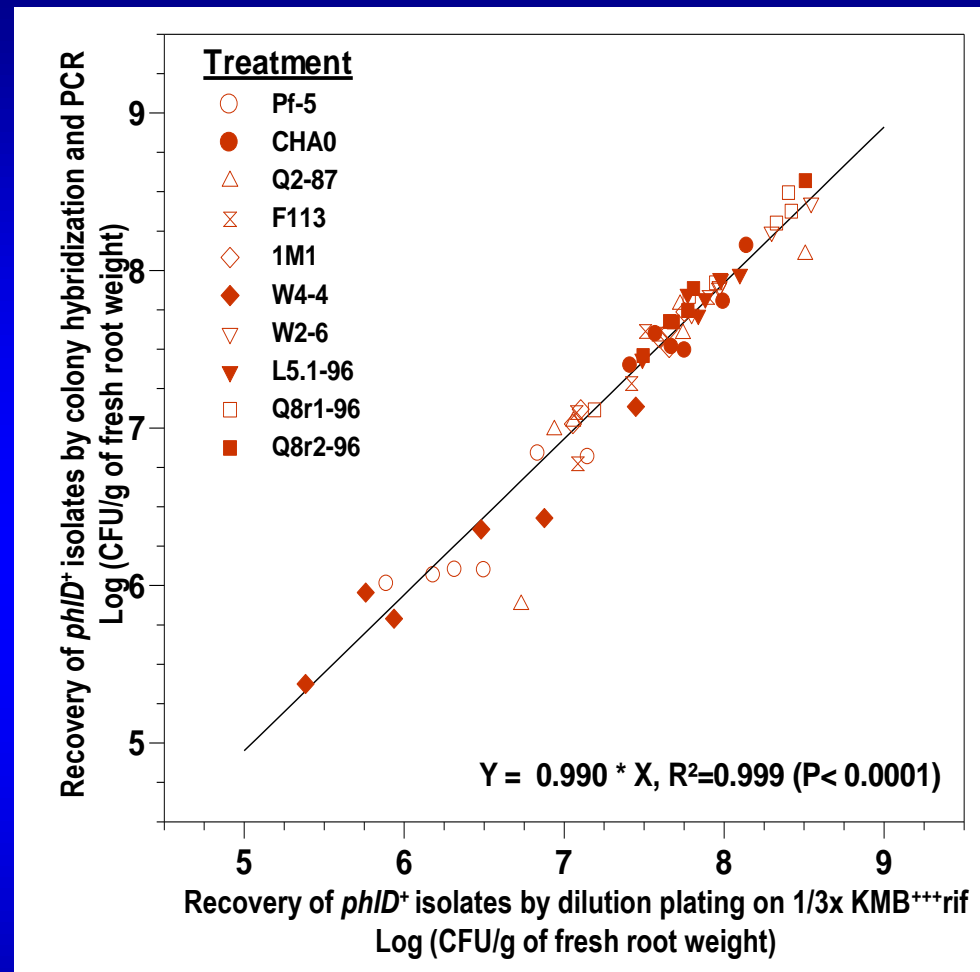
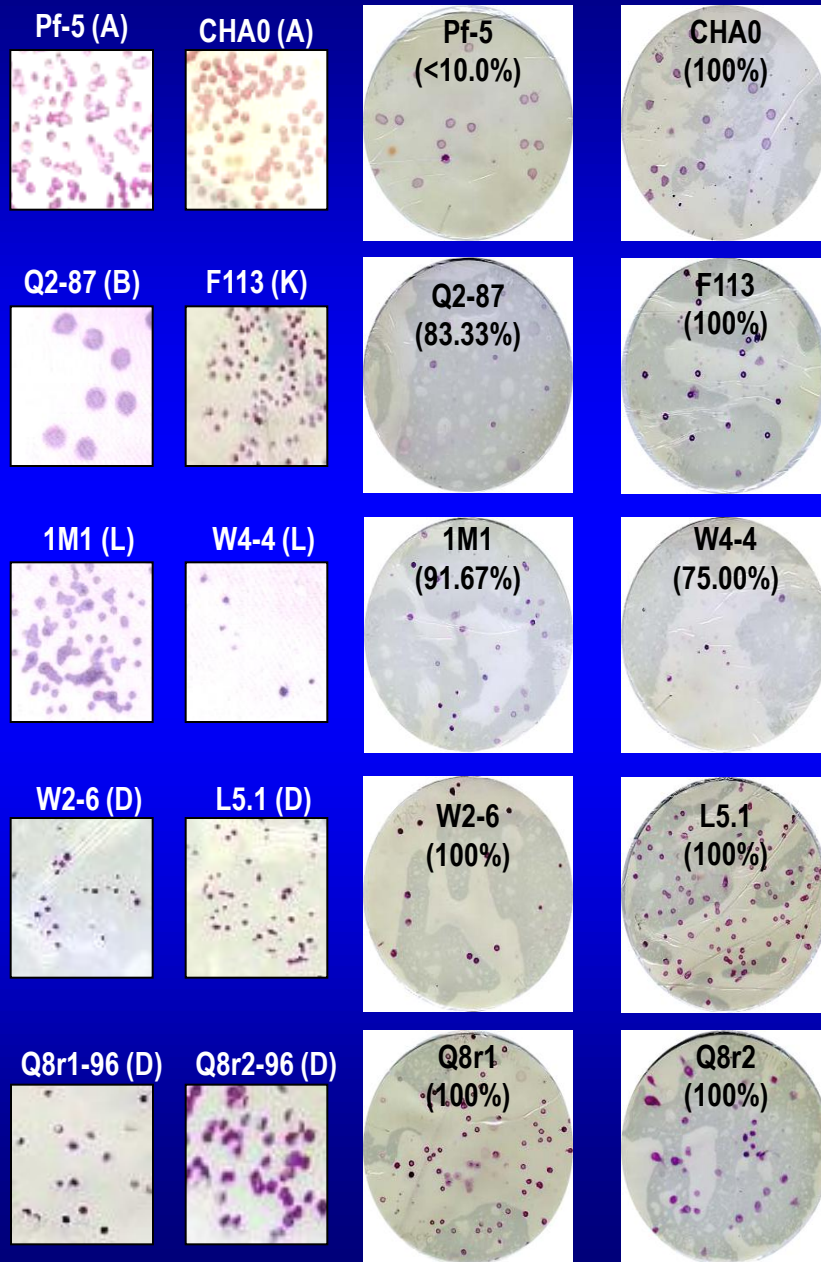
## PCR-based assay



## Plate dilution (1/3x KMB<sup>+++</sup>rif)



# Colony Hybridization & PCR vs. Antibiotic Resistance

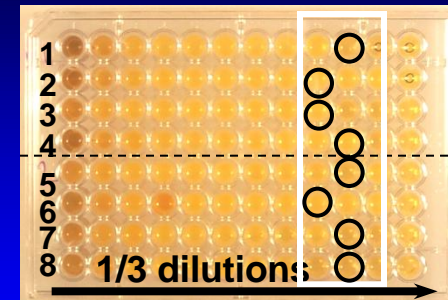


# Identification of DAPG-Producing *Pseudomonas* spp. from the Rhizosphere

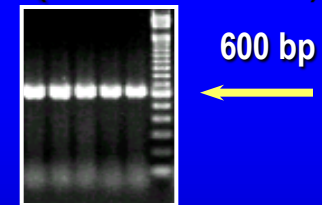
## Cycling experiments



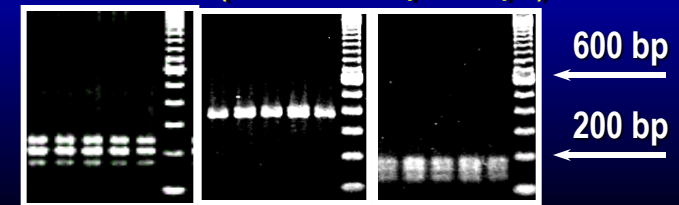
## PCR-based assay



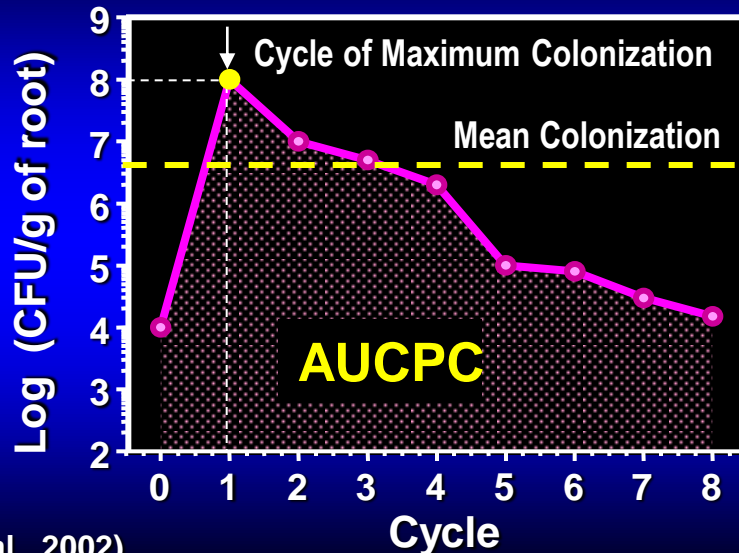
## PCR (B2BF/BPR4)



## RFLPs (*Hae*III, *Taq*I, *Msp*I)



## Comparison of treatments



(Landa et al., 2002)

(McSpadden Gardener et al., 2001)

# Tools for Tracking BCAs

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- Culture-Based Methods:

- ◆ PCR-Based Dilution-Endpoint Method:

- three fold dilution in microtiter plates with selective media followed by PCR with specific probes (McSpadden Gardener et al., 2001; Landa et al., 2002)

- Culture-Independent Quantitative PCR: (Rezzonico et al., 2003)

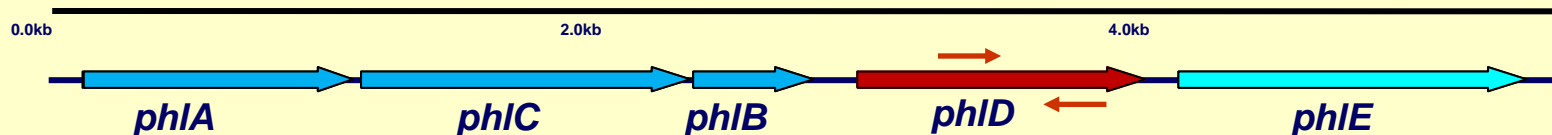
- Mavrodi et al., (2007) used a real-time PCR SYBR green assay to quantify different genotypes of DAPG-producing *P. fluorescens* on the roots of wheat.

# Genotype-specific *phlD* primers for real-time PCR

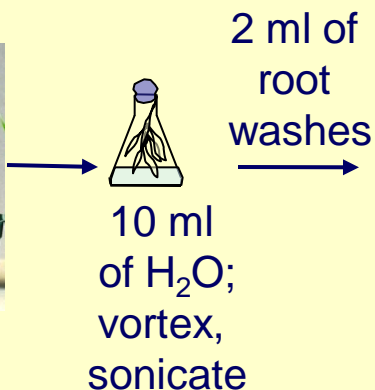
Genotype (strain)	Primers name	Primers sequence 5' to 3'	Positions	T <sub>m</sub> <sup>a</sup> of primers
<b>A</b> (Pf-5)	A_Up	GCACGGTGGAGGTTGGC	507	66.7 C
	A_Low	GTGATCGTCACTTCCTGCAC	575	62.6 C
<b>B</b> (Q2-87)	B_Up	CACGCATCCCAATTGAG	477	59.4 C
	B_Low	CCGTTACCTCTTGCACC	573	58.0 C
<b>D</b> (Q8r1-96)	D_Up	AGTTGCAGGACCAGTTC	37	57.6 C
<b>D</b> (FTAD1r34)	D_Low	CATTAAAGATGTCGCACCG	148	61.9 C
<b>I</b> (FTAD1r36)	I_Up	GGTTCCAGGTCCAGTTG	26	56.8 C
	I_Low	CGTCAAGGACAGTGGCTTC	199	62.9 C

<sup>a</sup> Oligonucleotides were designed by Oligo 6.65 Primer Analysis Software and T<sub>m</sub> of the primers were calculated by Oligo 6.65 using the nearest-neighbor thermodynamic method.

## *phl* biosynthesis genes



# General scheme for enumeration of introduced and indigenous bacteria in the wheat rhizosphere by real-time PCR



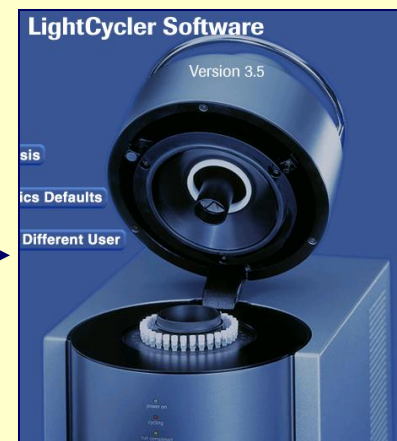
## UltraClean™ Soil DNA Isolation kit



Modified alternative protocol  
for wet soil samples

50 µL of DNA per sample

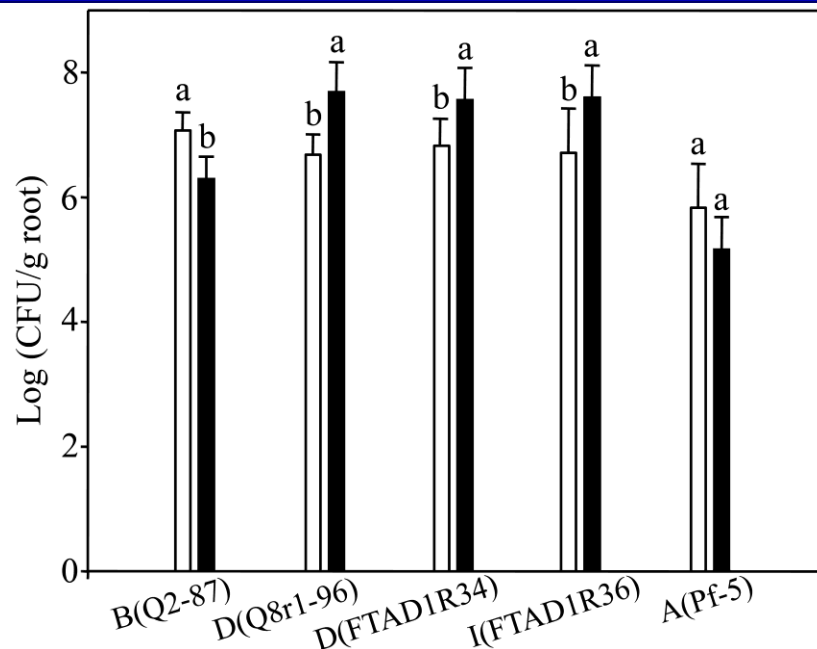
## Real-time PCR



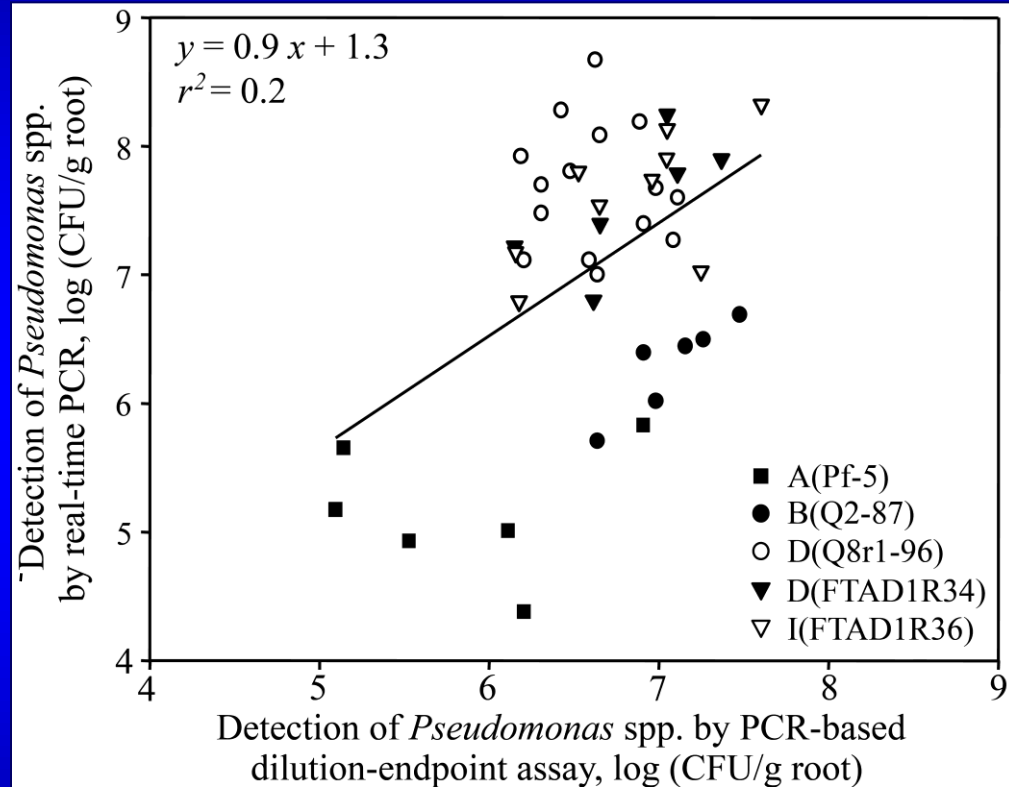
2 µL of DNA are  
used per PCR reaction



# Comparison of population levels of DAPG-producing *Pseudomonas* spp. in the rhizosphere of wheat detected by real-time PCR and TD endpoint assay



 Terminal dilution endpoint assay  
 real-time PCR



Regression analysis of population densities detected by real-time PCR and by a previously described TD endpoint assay indicated a significant linear relationship ( $P=0.0016$ ,  $r^2=0.2$ ). Each symbol represents the population of bacteria detected on roots plus adhering rhizosphere soil from a single seedling. [Mavrodi et al., 2007](#)

# Comparison of real-time PCR and TD endpoint assay

## Real-time PCR

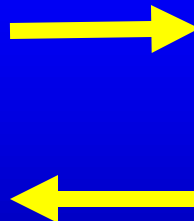
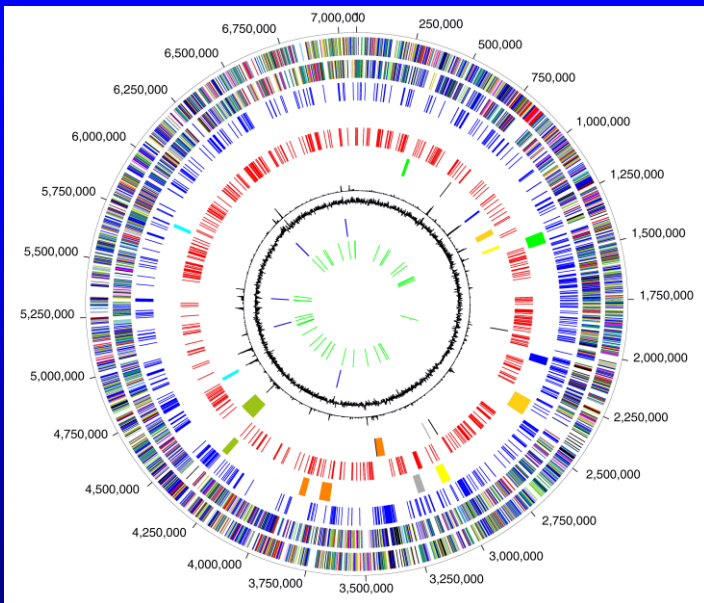
- Culture independent;
- Samples consist of purified DNA;
- Allows detection of dominant and subdominant genotypes;
- Allows detection of indigenous bacteria (i.e. those not tagged with antibiotic resistance);
- Detection limit on average is log 4 - 5 per rhizosphere;
- Cost per reaction ~ \$4.60 (including cost of soil DNA extraction, and not including cost of equipment and service contract);
- Turnaround time is 2 days.

## TD endpoint assay

- Culture dependent (inhibition may occur in mixed cultures during incubation step);
- Samples consist of frozen cell suspensions;
- Allows easy detection of only dominant genotypes;
- Works best for strains tagged with antibiotic resistance;
- Detection limit is log 3.3 CFU/g root;
- Cost per reaction ~ \$3.50;
- Turnaround time is 5 days.

# Thoughts for the Future

- Solving problems that are barriers to biocontrol reaching its full potential as an integral part of sustainable agriculture will require multi-disciplinary research focusing on the biocontrol process at all levels, from the genome to ecosystem scale.
- Developing tools to track the establishment and fate of BCAs will continue to be an important component of the multi-disciplinary research effort.







**Thank you**